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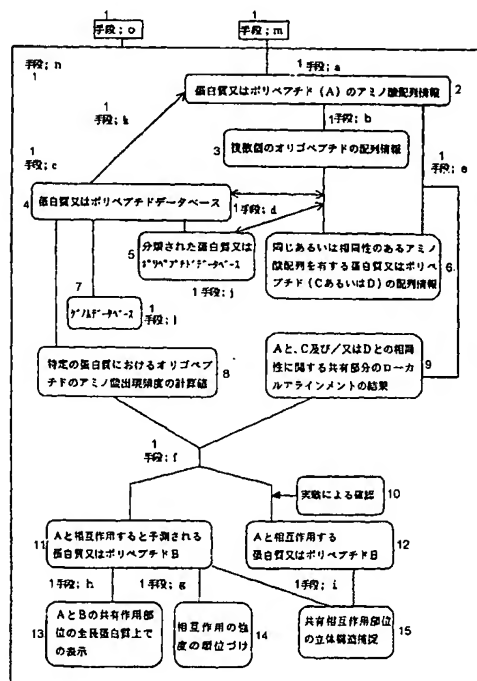
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(54) Title: METHOD OF ANTICIPATING INTERACTION BETWEEN PROTEINS

(54) 発明の名称: 蛋白質間相互作用予測方法



- 1... MEANS
- 2... AMINO ACID SEQUENCE DATA OF PROTEIN OR POLYPEPTIDE (A)
- 3... SEQUENCE DATA OF PLURAL OLIGOPEPTIDES
- 4... PROTEIN OR POLYPEPTIDE DATA BASE
- 5... CLASSIFIED PROTEIN OR POLYPEPTIDE DATA BASE
- 6... SEQUENCE DATA OF PROTEIN OR POLYPEPTIDE (C OR D) HAVING THE SAME OR HOMOLOGOUS AMINO ACID SEQUENCE
- 7... GENOME DATA BASE
- 8... CALCULATED AMINO ACID APPEARANCE FREQUENCIES OF OLIGOPEPTIDES IN SPECIFIC PROTEIN
- 9... RESULTS OF LOCAL ALIGNMENT OF PARTS COMMON TO A AND C AND/OR D CONCERNING HOMOLOGY
- 10... CONFIRMATION BY EXPERIMENT
- 11... PROTEIN OR POLYPEPTIDE B ANTICIPATED AS INTERACTING WITH A
- 12... PROTEIN OR POLYPEPTIDE B INTERACTING WITH A
- 13... INDICATION OF ACTION SITES COMMON TO A AND B ON FULL-LENGTH PROTEIN
- 14... INTERACTION STRENGTH RANKING
- 15... CERTIFICATION OF STERIC STRUCTURE OF COMMON INTERACTION SITES

(57) Abstract: A method of anticipating an interaction between proteins characterized by comprising: (1) digesting the amino acid sequence of a protein A to give oligopeptides of certain length; (2) searching for a protein C having the above-described oligopeptides or a protein D having oligopeptides homologous with the above-described oligopeptides from a protein data base; (3) performing local alignment between the above-described protein A and the protein C or D thus detected; and (4) anticipating that the detected protein C or D is

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